

01PE #2

Errors Corrected by the STIC Systems Branch

Serial Number: 09/844,915

CRF Processing Date: 5/11/2001
Entered by: AN (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

OIPE

RAW SEQUENCE LISTING

DATE: 05/11/2001

PATENT APPLICATION: US/09/844,915

TIME: 14:30:59

Input Set : A:\AP32737 sequence listing.txt

Output Set: N:\CRF3\05112001\I844915.raw

Does Not Comply
Corrected Diskette Needed

4 <110> APPLICANT Robbins, Paul D.
 5 Lu, Lina
 6 Giannoukakis, Nick
 8 <120> TITLE OF INVENTION: THE USE OF TOLEROGENIC DENDRITIC CELLS
 9 FOR ENHANCING TOLEROGENICITY IN A HOST AND METHODS FOR
 10 MAKING THE SAME
 12 <130> FILE REFERENCE: AP32737 / 072396.0225
 C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/844,915
 C--> 14 <141> CURRENT FILING DATE: 2001-04-27
 14 <150> PRIOR APPLICATION NUMBER: 60/200,479
 15 <151> PRIOR FILING DATE: 2000-04-28
 17 <160> NUMBER OF SEQ ID NOS: 7
 19 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

93 <210> SEQ ID NO: 7
 94 <211> LENGTH: 22
 95 <212> TYPE: DNA
 96 <213> ORGANISM: Artificial Sequence
 98 <220> FEATURE:
 99 <223> OTHER INFORMATION: Synthesized nucleotide sequence
 102 <400> SEQUENCE: 7
 103 agttgagggg actttccag gc
 E--> 110 ny02:322053.1

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VERIFICATION SUMMARY

DATE: 05/11/2001

PATENT APPLICATION: US/09/844,915

TIME: 14:31:00

Input Set : A:\AP32737 sequence listing.txt

Output Set: N:\CRF3\05112001\I844915.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:110 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:7
L:110 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:7
L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:110 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:26 SEQ:7
L:110 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
L:110 M:112 C: (48) String data converted to lower case,
L:110 M:252 E: No. of Seq. differs, <211>LENGTH:Input:22 Found:26 SEQ:7